




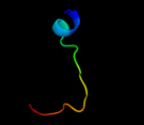


Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P0A8C8
Date	Thu Jan 5 11:07:28 GMT 2012
Unique Job ID	78563e4ea8665bbd

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3kw0D_	 Alignment		13.0	36	PDB header: hydrolase Chain: D: PDB Molecule: cysteine peptidase; PDBTitle: crystal structure of cysteine peptidase (np_982244.1) from bacillus2 cereus atcc 10987 at 2.50 a resolution
2	d1shyb2	 Alignment		8.8	42	Fold: Trefoil/Plexin domain-like Superfamily: Plexin repeat Family: Plexin repeat
3	c1q68A_	 Alignment		8.4	47	PDB header: membrane protein/transferase Chain: A: PDB Molecule: t-cell surface glycoprotein cd4; PDBTitle: solution structure of t-cell surface glycoprotein cd4 and2 proto-oncogene tyrosine-protein kinase lck fragments